INFORMATION DISCLOSURE CITATION *(Use several sheets if necessary)*	ATTY. DOCKET NO. PC25365A	SERIAL NO. 10/815,402
	APPLICANT Kimberly F. Fennell, et al.	
	FILING DATE March 31, 2004	GROUP 1614

OIPE NOV 2 3 2004 PATENT & TRADEMARK OFFICE

U.S. PATENT DOCUMENTS

EXAMINER INITIAL	DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE IF APPROPRIATE
NN	US 6 5 3 7 7 7 4	25 Mar03	Huang, et al.	453	69.1	

FOREIGN PATENT DOCUMENTS

	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION YES	TRANSLATION NO
NN	WO 0 3 4 8 7 3 3	12 Jun 03	International	G01N			

OTHER DOCUMENTS (*Including Author, Title, Date, Pertinent Pages, Etc.*)

NN	Robl, et al., U.S. Patent Application Publication No. US 2002/0035064, published March 21, 2002
	Kuntz, et al., "A Geometric Approach to Macromolecule-Ligand Interactions", *J. Mol. Biol.*, Vol. 161, pp. 269-288 (1982)
	Goodford, "A Computational Procedure for Determining Energetically Favorable Binding Sites on Biologically Important Macromolecules", *J. Med. Chem.*, Vol. 28, pp. 849-857 (1985)
	Goodsell, et al., "Automated Docking of Substrates to Proteins by Simulated Annealing", *Proteins: Structure, Function and Genetics*, Vol. 8, pp. 195-202 (1990)
	Miranker, et al., "Functionality Maps of Binding Sites: A Multiple Copy Simultaneous Search Method", *Proteins: Structure, Function and Genetics*, Vol. 11, pp. 29-34 (1991)
	Bartlett, et al., "Caveat: A Program to Facilitate the Structure-Derived Design of Biologically Active Molecules", *Molecular Recognition in Chemical and Biological Problems*, Spec. Pub. No. 78, Royal Chem. Soc., pp. 182-96 (1989)
	Martin, "3D Database Searching in Drug Design", *J. Med. Chem.*, Vol. 35, No. 12, pp. 2145-2154 (1992)
	Nishibata, et al., "Automatic Creation of Drug Candidate Structures Based on Receptor Structure. Starting Point for Artificial Lead Generation", *Tetrahedron*, Vol. 47, No. 43, pp. 8985-8990 (1991)
	Böhm, "The Computer Program LUDI: A new method for the de novo design of enzyme inhibitors", *J. Comp. Aid. Molec. Design*, Vol. 6 pp. 61-78 (1992)
	Cohen, et al., "Molecular Modeling Software and Methods for Medicinal Chemistry", *J. Med. Chem.*, Vol. 33, No. 3, pp. 883-894 (1990)
	Navia, et al., "Use of structural information in drug design", *Current Opinions in Structural Biology*, Vol. 2, pp. 202-210 (1992)
	Noren, et al. "A General Method for Site-Specific Incorporation of Unnatural Amino Acids Into Proteins", *Science*, Vol. 244, pp. 182-188 (1989)
	Fujihashi, et al., "Crystal Structure of *cis*-prenyl chain elongating enzyme, undecaprenyl diphosphate synthase", *PNAS*, Vol. 98, No. 8, pp. 4337-4342 (2001)
NN	Lattman, et al., "Use of the Rotation and Translation Functions", *Methods in Enzymology*, Vol. 115, pp. 55-77 (1985)

EXAMINER /Nashaat Nashed/ (06/20/2006) DATE CONSIDERED

EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.